

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
A0A0B4K813	A0A0B4K813_DROME	Drosophila melanogaster	NaN	25.496522	NaN	NaN	33925313	MCDSATTGCFLTRSSHRKENGSRVSP DVTASPGRAPPGLPANQMPAMGN QQHHGNQQHHGNQQHHGNQHS NHRGQSGSLSNAAGVKDPVMLQGD FRKVSIGSSEIFRQIEAVENDHDPNT AAALEAVERRGEMIVRVLEPRCMGS KQAVDAAHKLMNKADARHTVQLVE IVKRPQOTLGLYIREGNADRTDGVF ISRIALESAVYNSGCLRVGDEILAVN LVDVTHMSLDDVVIIMSIPRRLVLA RORRGNRGTGSPGPPTLSRPEQKPP PVVVIKRDLRDEDLDETRMPRPRS SRDRRTGDGREMTESRRLGLGLN NYSPPQEQDMMYNTNRGGGGGAMG EPPNWGYPKPPPPSSVITEOPTKAHA FAPSHAYYQNACTLESLEAEKVHAFY PGQPGGPPVGPSSRRMSTGTGNVGLA QQHARFPRSGSDQHLPRVEYSDYSN SLGRHSLLRSSLKPGTTGGAPMQVG VGGTLGRYGRYDQQRAGVSKYGPPS GGAQSLTRRSRPNLDYSSDTEATIGP RPSYYYYNRPAIGMSRGGGAGGG VGAASTAALLAGAADLNKFNLSLPRE RPGTRLQGIRSRMGDRLVDENDGN TSAPEFVRRRDLRQRITASPSIFT ADEYRAWLRRAPSSSAIAEQMRMTR DMFAQPRARFSCSAENIHDALRNT ESIYSSRNHILGTGLDRNMGLTRPI SALPVRSMSSQHIGGAGSIRSPSIRR MRQLELSAGPASPSGSILSTGGHQ SPAPTPSATLPRPHRQIDINPAEFAK YKLDKPIVDIGGISGMLWIHLLAGRG LRTAPEGAAGTATQGQTRDLYCVIEC DRVHKARTVVRSGDLQFDWDESEFE LDLVGNKQLDLVYVSWDPQHRHKL CYRGALSLSILRQSPHLQALKVEP RGTIYIRMRHTDPLALYKRRGLPSLR AGYPTLFGADLETVVNRESKNAPGS APVPIVLRRCVEEVERREGLDIIGLYRL CGSATKKRLLREAFERNRAVELTP EHVPDINVITGVLDKYLRELPEPLFT RCLFQMTVDALAVCLPDDPEGNAKL MLSILDCLPRANRATLVFLDHLSLV VNSERNKMSAQALATVMGPPLML HSASAPGADIDHAQPIAVLKYLLQI WPQQAQHQMAQHMGGAGAM MGGLVTAGSMSNMAGVASALSIGG LSGLNSRPIATATSRHTLPRQ	None	None	None	None	None	None	None		

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